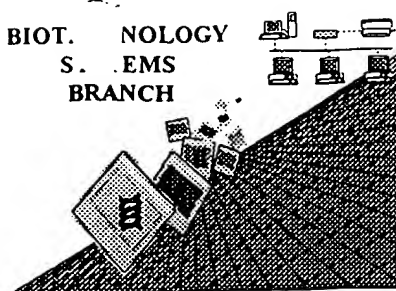


C. Collins

RAW SEQUENCE LISTING ERROR REPORT

BIOT. NOLOGY
S. EMS
BRANCH



#13

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/574,735

Source: 1600

Date Processed by STIC: 7-20-01

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

1600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/574,735

DATE: 07/20/2001

TIME: 13:18:29

Input Set : A:\cpg.txt

Output Set: N:\CRF3\07202001\I574735.raw

Does Not Comply
Corrected Diskette Needed
see p. 6

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3 <110> APPLICANT: De Veylder, Lieven
4     Beeckman, Tom
5     Inz,, Dirk
6     Van Camp, Wim
7     Krols, Luc
10 <120> TITLE OF INVENTION: Cyclin-dependent kinase inhibitors and uses thereof
13 <130> FILE REFERENCE: 2283/301
16 <140> CURRENT APPLICATION NUMBER: US 09/574,735
17 <141> CURRENT FILING DATE: 2000-05-18
20 <160> NUMBER OF SEQ ID NOS: 48
23 <170> SOFTWARE: PatentIn version 3.0
26 <210> SEQ ID NO: 1
27 <211> LENGTH: 932
28 <212> TYPE: DNA
29 <213> ORGANISM: Arabidopsis thaliana
31 <220> FEATURE:
32 <221> NAME/KEY: CDS
33 <222> LOCATION: (86)..(712)
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37 ctggaagggtg acgtcgtagg agatt atg gcg gcg gtt agg aga aga gaa cga      112
38                               Met Ala Ala Val Arg Arg Arg Glu Arg
39                               1           5
40 gat gtg gtt gaa gag aat gga gtt acg acg acg acg gtg aaa cga agg      160
41 Asp Val Val Glu Glu Asn Gly Val Thr Thr Thr Thr Val Lys Arg Arg
42 10                               15           20           25
43 aag atg gag gag gaa gtg gat tta gtg gaa tct agg ata att ctg tct      208
44 Lys Met Glu Glu Glu Val Asp Leu Val Glu Ser Arg Ile Ile Leu Ser
45                               30           35           40
46 ccg tgt gta cag gcg acg aat cgc ggt gga att gtg gcg aga aat tca      256
47 Pro Cys Val Gln Ala Thr Asn Arg Gly Gly Ile Val Ala Arg Asn Ser
48                               45           50           55
49 gca gga gcg tcg gag acg agt gtt gtt ata gta cga cgg cga gat tct      304
50 Ala Gly Ala Ser Glu Thr Ser Val Val Ile Val Arg Arg Arg Asp Ser
51                               60           65           70
52 cct ccg gtt gaa gaa cag tgt caa atc gaa gaa gaa gat tcg tcg gtt      352
53 Pro Pro Val Glu Glu Gln Cys Gln Ile Glu Glu Glu Asp Ser Ser Val
54                               75           80           85
55 tcg tgt tgt tct aca tcg gaa gag aaa tcg aaa cgg aga atc gaa ttt      400
56 Ser Cys Cys Ser Thr Ser Glu Glu Lys Ser Lys Arg Arg Ile Glu Phe
57 90                               95           100          105
58 gta gat ctt gag gaa aat aac ggt gac gat cgt gaa aca gaa acg tcg      448
59 Val Asp Leu Glu Glu Asn Asn Gly Asp Asp Arg Glu Thr Glu Thr Ser
60                               110          115          120
61 tgg att tac gat gat ttg aat aag agt gag gaa tcg atg aac atg gat      496
62 Trp Ile Tyr Asp Asp Leu Asn Lys Ser Glu Glu Ser Met Asn Met Asp
63                               125          130          135

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RAW SEQUENCE LISTING

DATE: 07/20/2001

PATENT APPLICATION: US/09/574,735

TIME: 13:18:29

Input Set : A:\cpg.txt

Output Set: N:\CRF3\07202001\I574735.raw

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64 tct tct tcg gtg gct gtt gaa gat gta gag tct cgc cgc agg tta agg      544
65 Ser Ser Ser Val Ala Val Glu Asp Val Glu Ser Arg Arg Arg Leu Arg
66          140          145          150
67 aag agt ctc cat gag acg gtg aag gaa gct gag tta gaa gat ttt ttt      592
68 Lys Ser Leu His Glu Thr Val Lys Glu Ala Glu Leu Glu Asp Phe Phe
69          155          160          165
70 cag gtg gcg gag aaa gat ctt cgg aat aag ttg ttg gaa tgt tct atg      640
71 Gln Val Ala Glu Lys Asp Leu Arg Asn Lys Leu Leu Glu Cys Ser Met
72 170          175          180          185
73 aag tat aac ttc gat ttc gag aaa gat gag cca ctt ggt gga gga aga      688
74 Lys Tyr Asn Phe Asp Phe Glu Lys Asp Glu Pro Leu Gly Gly Gly Arg
75          190          195          200
76 tac gag tgg gtt aaa ttg aat cca tgaagaagac gatgatgata atgatgatca      742
77 Tyr Glu Trp Val Lys Leu Asn Pro
78          205
79 ttgtttttcac caaagtactt attatttttt tttctgtaata atcttttgctt tgattttttct      802
80 ttttaacaaaa tccaaatgta gatatctttt tctcgaataa tcaataacat gtaattcaac      862
81 ttttgtttgt acttccttga ggtaattaat tagattcgtg tttttctcga ttaataaaact      922
82 ataagtttat      932
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86 <211> LENGTH: 209
87 <212> TYPE: PRT
88 <213> ORGANISM: Arabidopsis thaliana
90 <400> SEQUENCE: 2
91 Met Ala Ala Val Arg Arg Arg Glu Arg Asp Val Val Glu Glu Asn Gly
92 1          5          10          15
93 Val Thr Thr Thr Thr Val Lys Arg Arg Lys Met Glu Glu Glu Val Asp
94          20          25          30
95 Leu Val Glu Ser Arg Ile Ile Leu Ser Pro Cys Val Gln Ala Thr Asn
96          35          40          45
97 Arg Gly Gly Ile Val Ala Arg Asn Ser Ala Gly Ala Ser Glu Thr Ser
98          50          55          60
99 Val Val Ile Val Arg Arg Arg Asp Ser Pro Pro Val Glu Glu Gln Cys
100 65          70          75          80
101 Gln Ile Glu Glu Glu Asp Ser Ser Val Ser Cys Cys Ser Thr Ser Glu
102          85          90          95
103 Glu Lys Ser Lys Arg Arg Ile Glu Phe Val Asp Leu Glu Glu Asn Asn
104          100          105          110
105 Gly Asp Asp Arg Glu Thr Glu Thr Ser Trp Ile Tyr Asp Asp Leu Asn
106          115          120          125
107 Lys Ser Glu Glu Ser Met Asn Met Asp Ser Ser Ser Val Ala Val Glu
108          130          135          140
109 Asp Val Glu Ser Arg Arg Arg Leu Arg Lys Ser Leu His Glu Thr Val
110 145          150          155          160
111 Lys Glu Ala Glu Leu Glu Asp Phe Phe Gln Val Ala Glu Lys Asp Leu
112          165          170          175
113 Arg Asn Lys Leu Leu Glu Cys Ser Met Lys Tyr Asn Phe Asp Phe Glu
114          180          185          190
115 Lys Asp Glu Pro Leu Gly Gly Gly Arg Tyr Glu Trp Val Lys Leu Asn

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/574,735

DATE: 07/20/2001

TIME: 13:18:29

Input Set : A:\cpg.txt

Output Set: N:\CRF3\07202001\I574735.raw

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117 Pro
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122 <211> LENGTH: 875
123 <212> TYPE: DNA
124 <213> ORGANISM: Arabidopsis thaliana
126 <220> FEATURE:
127 <221> NAME/KEY: CDS
128 <222> LOCATION: (11)..(658)
130 <400> SEQUENCE: 3
131 ggcacgagag aaa tca aag ata act ggc gat atc agc gtc atg gaa gtc      49
132          Lys Ser Lys Ile Thr Gly Asp Ile Ser Val Met Glu Val
133          1          5          10
134 tct aaa gca aca gct cca agt cca ggt gtt cga acc aga gcc gct aaa      97
135 Ser Lys Ala Thr Ala Pro Ser Pro Gly Val Arg Thr Arg Ala Ala Lys
136          15          20          25
137 acc cta gcc ttg aag cgg ctt aat tcc tcc gcc gct gat tca gct cta      145
138 Thr Leu Ala Leu Lys Arg Leu Asn Ser Ser Ala Ala Asp Ser Ala Leu
139 30          35          40          45
140 cct aac gac tct tct tgc tat ctt cag ctc cgt agc cgc cgt ctc gag      193
141 Pro Asn Asp Ser Ser Cys Tyr Leu Gln Leu Arg Ser Arg Arg Leu Glu
142          50          55          60
143 aaa ccc tct tgc ctg att gaa ccg aaa cag ccg ccg aga gtt cac aga      241
144 Lys Pro Ser Ser Leu Ile Glu Pro Lys Gln Pro Pro Arg Val His Arg
145          65          70          75
146 tcg gga att aaa gag tct ggt tcc agg tct cgc gtt gac tcg gtt aac      289
147 Ser Gly Ile Lys Glu Ser Gly Ser Arg Ser Arg Val Asp Ser Val Asn
148          80          85          90
149 tcg gtt cct gta gct cag agc tct aat gaa gat gaa tgt ttt gac aat      337
150 Ser Val Pro Val Ala Gln Ser Ser Asn Glu Asp Glu Cys Phe Asp Asn
151          95          100          105
152 ttc gtg agt gtc caa gtt tct tgt ggt gaa aac agt ctc ggt ttt gaa      385
153 Phe Val Ser Val Gln Val Ser Cys Gly Glu Asn Ser Leu Gly Phe Glu
154 110          115          120          125
155 tca aga cac agc aca agg gag agc acg cct tgt aac ttt gtt gag gat      433
156 Ser Arg His Ser Thr Arg Glu Ser Thr Pro Cys Asn Phe Val Glu Asp
157          130          135          140
158 atg gag atc atg gtt aca cca ggg tct agc acg agg tcg atg tgc aga      481
159 Met Glu Ile Met Val Thr Pro Gly Ser Ser Thr Arg Ser Met Cys Arg
160          145          150          155
161 gca acc aaa gag tac aca agg gaa caa gat aac gtg atc ccg acc act      529
162 Ala Thr Lys Glu Tyr Thr Arg Glu Gln Asp Asn Val Ile Pro Thr Thr
163          160          165          170
164 agt gaa atg gag gag ttc ttt gca tat gca gag cag cag caa cag agg      577
165 Ser Glu Met Glu Glu Phe Phe Ala Tyr Ala Glu Gln Gln Gln Gln Arg
166          175          180          185
167 cta ttc atg gag aag tac aac ttc gac att gtg aat gat atc ccc ctc      625
168 Leu Phe Met Glu Lys Tyr Asn Phe Asp Ile Val Asn Asp Ile Pro Leu
169 190          195          200          205

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RAW SEQUENCE LISTING

DATE: 07/20/2001

PATENT APPLICATION: US/09/574,735

TIME: 13:18:29

Input Set : A:\cpg.txt

Output Set: N:\CRF3\07202001\I574735.raw

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170 agc gga cgt tac gaa tgg gtg caa gtc aaa cca tgaagttcaa aaggaaacag 678
171 Ser Gly Arg Tyr Glu Trp Val Gln Val Lys Pro
172          210          215
173 ctccaaaaga catggtgtga agttagagaa tgtgatggag ttaacagact aaccaaacat 738
174 cagaaatcgt gtaatcttaa gtaataatgt ggtagagaa caagtttgag agtagcttag 798
175 ggaccttaaa acctcacacc atttgtaata ctaatcttct tcagatgctt agtgaaattt 858
176 tctcatctgt ttctttc 875
179 <210> SEQ ID NO: 4
180 <211> LENGTH: 216
181 <212> TYPE: PRT
182 <213> ORGANISM: Arabidopsis thaliana
184 <400> SEQUENCE: 4
185 Lys Ser Lys Ile Thr Gly Asp Ile Ser Val Met Glu Val Ser Lys Ala
186 1          5          10          15
187 Thr Ala Pro Ser Pro Gly Val Arg Thr Arg Ala Ala Lys Thr Leu Ala
188          20          25          30
189 Leu Lys Arg Leu Asn Ser Ser Ala Ala Asp Ser Ala Leu Pro Asn Asp
190          35          40          45
191 Ser Ser Cys Tyr Leu Gln Leu Arg Ser Arg Arg Leu Glu Lys Pro Ser
192          50          55          60
193 Ser Leu Ile Glu Pro Lys Gln Pro Pro Arg Val His Arg Ser Gly Ile
194 65          70          75          80
195 Lys Glu Ser Gly Ser Arg Ser Arg Val Asp Ser Val Asn Ser Val Pro
196          85          90          95
197 Val Ala Gln Ser Ser Asn Glu Asp Glu Cys Phe Asp Asn Phe Val Ser
198          100         105         110
199 Val Gln Val Ser Cys Gly Glu Asn Ser Leu Gly Phe Glu Ser Arg His
200          115         120         125
201 Ser Thr Arg Glu Ser Thr Pro Cys Asn Phe Val Glu Asp Met Glu Ile
202          130         135         140
203 Met Val Thr Pro Gly Ser Ser Thr Arg Ser Met Cys Arg Ala Thr Lys
204 145          150         155         160
205 Glu Tyr Thr Arg Glu Gln Asp Asn Val Ile Pro Thr Thr Ser Glu Met
206          165         170         175
207 Glu Glu Phe Phe Ala Tyr Ala Glu Gln Gln Gln Gln Arg Leu Phe Met
208          180         185         190
209 Glu Lys Tyr Asn Phe Asp Ile Val Asn Asp Ile Pro Leu Ser Gly Arg
210          195         200         205
211 Tyr Glu Trp Val Gln Val Lys Pro
212          210         215
215 <210> SEQ ID NO: 5
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217 <212> TYPE: DNA
218 <213> ORGANISM: Arabidopsis thaliana
220 <220> FEATURE:
221 <221> NAME/KEY: CDS
222 <222> LOCATION: (92)..(763)
224 <400> SEQUENCE: 5
225 aaaccactct tcaaatcaaa cactttctta cataagattc ctctgttttt ctgtgtgctt 60

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/574,735

DATE: 07/20/2001

TIME: 13:18:29

Input Set : A:\cpg.txt

Output Set: N:\CRF3\07202001\I574735.raw

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226 cttcaaattc ttccctgtt tttcaacttc a / atg ggg aag tac atg aag aaa      112
227                               Met Gly Lys Tyr Met Lys Lys
228                               1                               5
229 ctc aaa tcc aaa tca gaa tct cct tca ccc aat tca aca cca aca cca      160
230 Leu Lys Ser Lys Ser Glu Ser Pro Ser Pro Asn Ser Thr Pro Thr Pro
231          10          15          20
232 tca cca tca cca tca cca aca cca atc acc acc aat tca cca cca cca      208
233 Ser Pro Ser Pro Ser Pro Thr Pro Ile Thr Thr Asn Ser Pro Pro Pro
234          25          30          35
235 aca aca ccc aat tcc tct gat ggt gtt cga act cgt gct aga acc cta      256
236 Thr Thr Pro Asn Ser Ser Asp Gly Val Arg Thr Arg Ala Arg Thr Leu
237 40          45          50          55
238 gct ttg gag aat tcc aac aat cag aat cag aat ctt tct gtt tct tct      304
239 Ala Leu Glu Asn Ser Asn Asn Gln Asn Gln Asn Leu Ser Val Ser Ser
240          60          65          70
241 gat tct tac ctt cag ctg agg aac cgt cgc ctt aag aga ccc cta att      352
242 Asp Ser Tyr Leu Gln Leu Arg Asn Arg Arg Leu Lys Arg Pro Leu Ile
243          75          80          85
244 agg caa cat tcc gct aag agg aat aag ggg cat gat gga aac cct aaa      400
245 Arg Gln His Ser Ala Lys Arg Asn Lys Gly His Asp Gly Asn Pro Lys
246          90          95          100
247 tcc cca att ggg gat tca att gct gaa gag aaa act gtt cag aag agt      448
248 Ser Pro Ile Gly Asp Ser Ile Ala Glu Glu Lys Thr Val Gln Lys Ser
249          105          110          115
250 cct gag cct gaa aat gct gaa ttc aag gag aat gct gag gat act gag      496
251 Pro Glu Pro Glu Asn Ala Glu Phe Lys Glu Asn Ala Glu Asp Thr Glu
252 120          125          130          135
253 aga agc gct agg gaa act aca ccc gtc cat ttg ata atg cga gca gac      544
254 Arg Ser Ala Arg Glu Thr Thr Pro Val His Leu Ile Met Arg Ala Asp
255          140          145          150
256 gtt ctc agg cct cct agg cca att acc agg cgt act ttt cca act gaa      592
257 Val Leu Arg Pro Pro Arg Pro Ile Thr Arg Arg Thr Phe Pro Thr Glu
258          155          160          165
259 gct aat ccc aaa acg gag cag cca act atc cca att tca cgc gaa ttt      640
260 Ala Asn Pro Lys Thr Glu Gln Pro Thr Ile Pro Ile Ser Arg Glu Phe
261          170          175          180
262 gag gaa ttc tgt gct aaa cat gaa gcc gag cag caa agg gag ttc atg      688
263 Glu Glu Phe Cys Ala Lys His Glu Ala Glu Gln Gln Arg Glu Phe Met
264          185          190          195
265 gag aag tac aac ttt gat cct gtg aca gag cag cca ctc cca ggg cgt      736
266 Glu Lys Tyr Asn Phe Asp Pro Val Thr Glu Gln Pro Leu Pro Gly Arg
267 200          205          210          215
268 tac gaa tgg gaa aaa gtg tcg ccc tag aaggcaggct agtattaagt      783
269 Tyr Glu Trp Glu Lys Val Ser Pro
270          220
271 gttccatcaa tacatcttta aagtagcagc agggtagaa tttgttgaaa aggggtggtgg      843
272 tgctatttcc attttccatc actttctatt tacttgtaaa gaaagtagga ctttcaacat      903
273 atgtagacta atgatctgta actttacaga ggtgttgatt acacaacaat acaaagtcct      963
274 ttgtctagca gatcattaaa gaagggtttg agggaataag ggtctctagt tgtaggggtt      1023

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/574,735

DATE: 07/20/2001

TIME: 13:18:30

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Output Set: N:\CRF3\07202001\I574735.raw

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/574,735

DATE: 07/20/2001

TIME: 13:18:30

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L:615 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36
L:629 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37
L:641 M:283 W: Missing Blank Line separator, <220> field identifier
L:644 M:283 W: Missing Blank Line separator, <220> field identifier
L:649 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38
L:663 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39